rs Corrected by the STIC System Sranch CRF Processing Date: Serlal Number: Edited by: Verified by: Changed a file from non-ASCII to ASCII Changed the margins in cases where the sequence text was "wrapped" down to the next line. Edited a format error in the Current Application Data section, specifically Edited the Current Application Data section with the actual current number. The number inputted by the applicant was 🔲 the prior application data; or 🔲 other Added the mandatory heading and subheadings for "Current Application Data". Edited the "Number of Sequences" field. The applicant spelled out a number instead of using an integer. Changed the spelling of a mandatory field (the headings or subheadings), specifically: Corrected the SEQ ID NO when obviously incorrect. The sequence numbers that were edited were: Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited: Corrected subheading placement. All responses must be on the same line as each subheading. If the applicant placed a response below the subheading, this was moved to its appropriate place. Inserted colons after headings/subheadings. Headings edited included: Deleted extra, invalid, headings used by an applicant, specifically: Deleted: non-ASCII "garbage" at the beginning/end of files; secretary initials/filename at end of file; page numbers throughout text; other invalid text, such as\_\_\_\_\_ Inserted mandatory headings, specifically: Corrected an obvious error in the response, specifically: Edited identifiers where upper case is used but lower case is required, or vice versa. Corrected an error in the Number of Sequences field, specifically: A "Hard Page Break" code was inserted by the applicant. All occurrences had to be deleted. Deleted ending stop codon in amino acid sequences and adjusted the "(A)Length:" field accordingly (error due to a Patentin bug). Sequences corrected: \_\_\_\_\_ Other:

\*Examiner: The above corrections must be communicated to the applicant in the first Office Action. DO NOT send a copy of this form.





1600

RAW SEQUENCE LISTING DATE: 05/06/2002 FATENT APPLICATION: US/09/928,0478 TIME: 17:55:53

Input Set : N:\jumbos\928047B.txt

Cutput Set: N:\CRF3\05062002\1928047B.raw

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4 -1105 AFFLICANT: Cantor, Thomas
 5 KILON TITLE OF INVENTION: CYCLASE INHIBITING PARATHYROID HORMONE
         ANTAGONIST OR MODULATORS AND OSTEOPOROSIS
10 <1300 FILE FEFERENCE: 53221-20002.00
12 <140> CURRENT APPLICATION NUMBER: US 09/928,047B
15 (141> CUFRENT FILING DATE: 2001-08-10
15 <150> PRIOR APPLICATION NUMBER: US 60/221,440
16 <151> PRIOR FILING DATE: 2000-08-13
18 <160> NUMBER OF SEQ ID NOS: 8
20 <170% SOFTWARE: FastSEQ for Windows Version 4.0
02 <210% SEQ ID NO: 1
13 <2015 LENGTH: 83
24 <212> TYPE: PRT
25 «Il3» ORGANISM: Homo sapiens
17 <400 - SEQUENCE: 1
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30 Met Glu Arg Val Glu Trp Leu Arg Lys Lys Leu Gln Asp Val His Asn
                                    2.5
               ž C
: Phe Val Ala Leu Gly Ala Pro Leu Ala Pro Arg Asp Ala Gly Ser Gln
                               4.1
          3 5.
.5 .5
34 Ard Pro Arg Lys Lys Gld Asp Asn Val Leu Val Glu Ser His Glu Lys
                           5.5
                                                \epsilon 0
Fr Ser Let Gly Gld Ala Ash Lys Ala Asp Val Ash Val Leu Thr Lys Ala
311 65
38 Lys Ser Gln
41 <710× SEQ ID NO: 2
41 4111 - LENGTH: 52
43 KILL TYPE: PRT
44 K213~ ORGANISM: Homo sapiens
4r <4 0% SEQUENCE: 2
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4 4
                   Ε.
                                        10
40 Glo Arg Val Glu Trp Leu Arg Lys Lys Leu Gln Asp Val His Asn Phe
                                    2:5
               20
51 Val Ala Leu Bly Ala Pro Leu Ala Pro Arg Asp Ala Gly Ser Gln Arg
                                40
to Pro Arg Lys Lys Glu Asp Asn Val Leu Val Glu Ser His Glu Lys Ser
                            5, 5,
  Let. Gly Glu Ala Asn Lys Ala Asp Val Asn Val Leu Thr Lys Ala Lys
54 65
5° Ser Gln
67 (210> SEQ ID NO: 3
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RAW SEQUENCE LISTING DATE: 05/06/2002 PATENT APPLICATION: US/09/928,047B TIME: 17:55:53

Input Set : N:\jumbos\928047B.txt

Output Set: N:\CRF3\05062002\1928047B.raw

81 4214> LENGTH: 51 50 HU12> TYPE: PRT 53 Kuli3> DRGANIŠM: Homo sapiens 6% 14005 SEQUENCE: 3 So Phe Val Ala Leu Gly Ala Pro Leu Ala Pro Arg Asp Ala Gly Ser Gln 1 5, 10 65 Arg Pro Arg Lys Lys Glu Asp Asn Val Leu Val Glu Ser His Glu Lys 2.5 5.4 70 Ser Leu Gly Glu Ala Ash Lys Ala Asp Val Ash Val Leu Thr Lys Ala 3.5 4.0 71 Lys Ser Gln 5.5 75 %1100 SEQ ID NO: 4 76 <5110 LENGTH: 78  $7^{\prime\prime\prime}$  <112> TYPE: PFT 78 (113) ORGANISM: Homo sapiens 80 K400> SEQUENCE: 4 31 Leu Met His Asn Leu Gly Lys His Leu Asn Ser Met Glu Arg Val Glu 1.0 3) Trp Leu Arg Lys Lys Leu Gln Asp Val His Asn Phe Val Ala Leu Gly 25 26 85 Ala Pro Leu Ala Pro Arg Asp Ala Gly Ser Gln Arg Pro Arg Lys Lys 4.5 3.5 4.0 87 Glu Asp Ash Val Leu Val Glu Ser His Glu Lys Ser Leu Gly Glu Ala Ę, O 55 80 89 Ash Lys Ala Asp Val Ash Val Leu Thr Lys Ala Lys Ser Gln 90 65 70 91 <110> SEQ ID NO: 5 9 - <1115 LENGTH: 84 94 (L125 TYPE: PRT 35 (213) ORGANISM: Homo sapiens #1 k4+0> SEQUENCE: 5 Pr Val Ser Glu Ile Gln Leu Met His Asn Leu Gly Lys His Leu Asn 1.0 100 Ser Met Glu Arg Val Glu Trp Leu Arg Lys Lys Leu Gln Asp Val His 25 101 i C 1'2 Asn Phe Val Ala Lei Gly Ala Pro Leu Ala Pro Arg Asp Ala Gly Ser 1 (3) 35 4 0 14 Gln Arg Fro Arg Lys Lys Glu Asp Asn Val Leu Val Glu Ser His Glu 115 50 55 106 Lys Ser Leu Gly Glu Ala Ash Lys Ala Asp Val Ash Val Leu Thr Lys 107 65 80 7.0 1 8 Ala Lys Ser Gln 1.1 (1105 SEQ ID NO: 6 112 <.115 LENGTH: 34 113 <212> TYPE: PRT 114 (213) OF GANISM: Homo sapiens 116 4400> SEQUENCE: 6

117 Ser Val Ser Glu Ile Gln Leu Met His Ash Leu Gly Lys His Leu Ash

RAW SEQUENCE LISTING

FATENT APPLICATION: US/09/928,047B

DATE: 05/08/2002 TIME: 17:55:53

Input Set : N:\jumbos\928047B.txt

Output Set: N:\CRF3\05062002\I928047B.raw

118 1 119 Ser Met Glu Arg Val Glu Trp Leu Arg Lys Lys Leu Gln Asp Val His 111 25 ld. Asn Phe 184 K010> SEQ ID NO: 7 1.3 <1211> LENGTH: 50 114 #112> TYPE: PRT 1.1 (113> ORGANISM: Homo sapiens 11 % %400> SEQUENCE: 7 15: Val Ala Leu Gly Ala Pro Leu Ala Pro Arg Asp Ala Gly Ser Gln Arg Ξ, 10 13. Pro Arg Lys Lys Glu Asp Asn Val Leu Val Glu Ser His Glu Lys Ser 135 25 134 Leu Gly Glu Ala Asn Lys Ala Asp Val Asn Val Leu Thr Lys Ala Lys 135 35 40 138 Ser Glr 50 137 13 + <2:10> SEQ ID NO: 8 14 + <211> LENGTH: 57 141 <212> TYPE: PRT 142 <213> ORGANISM: Homo sapiens 144 <400> SEQUENCE: 8 145 Leu Gln Asp Val His Asn Phe Val Ala Leu Gly Ala Pro Leu Ala Pro 146 1 5 10 14: Arg Asp Ala Gly Ser Gln Arg Pro Arg Lys Lys Glu Asp Asn Val Leu 2.6 25 14 -140 Val Glu Ser His Glu Lys Ser Leu Gly Glu Ala Asn Lys Ala Asp Val 170 35 4.0 15: Asn Val Leu Thr Lys Ala Lys Ser Gln

152 50

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/928,047B

DATE: 05/06/2002 TIME: 17:55:54

Input Set : N:\jumbos\928047B.txt

Output Set: N:\CRF3\05062002\I928047B.raw